

## NEW JERSEY LAB ID#:20012 : NEW YORK LAB ID#: 11376

## GC/MS SEMI-VOLATILE ANALYSIS CONFORMANCE/NON-CONFORMANCE SUMMARY

CHEMTECH PROJEC	CT NUMBER: BP041725			
SequenceID :	BP041725	NA	NO	YES
1. Chromatograms Lal	peled/Compounds Identified. (Field samples and Method Blank	(3)		✓
2. GC/MS Tuning Specifications. DFTPP Meet Criteria Criteria (NOTE THAT THERE ARE DIFFERENT CRITERIA FOR NY ASP CLP, CLP AND NJ)				<b>_</b>
3. GC/MS Tuning Free series	quency - Performed every 24 hours for 600 series and 12 hours	for 8000		<b>√</b>
analysis and continu	- Initial Calibration performed within 30 days before sample sing calibration performed within 24 hours of sample analysis 2 hours for 8000 series			✓
5. GC/MS Calibration	Met:			✓
a. Initial calibration If not met, list those co	Meet Criteria ompounds and their recoveries which fall outside the acceptable	range.		✓
If not met, list those co	ration(CCC) Meet Criteria ompounds and their recoveries which fall outside the acceptable	e range.	<u> </u>	
6. Blank Contaminatio	arginally biased high in the CCC. on - If yes, list compounds and concentrations in each blank:		_✓	
a. B/N Fraction				

d. Acid Fraction

<ul> <li>7. Surrogate Recoveries Meet Criteria</li> <li>If not met, list those compounds and their recoveries which fall outside the acceptable ranges.</li> <li>a. B/N Fraction</li> </ul>		✓	
<ul> <li>d. Acid Fraction</li> <li>8. Matrix Spike/Matrix Spike Duplicate Recoveries Meet Criteria</li> <li>If not met, list those compounds and their recoveries which fall outside the acceptable range.</li> <li>a. B/N Fraction Recovery failed for some compound in Q1821-01MS/MSD due to matrix interference.</li> </ul>		<b>√</b>	
<ul> <li>d. Acid Fraction</li> <li>9. Internal Standard Area/Retention Time Shift Meet Criteria</li> </ul>		<u></u>	
Comments: Internal standard failed in Q1825-01, Q1812-03, Q1826-03. these samples will be re-analyzed to confirm the matu interference.	rix		
10. Extraction Holding Time Met         If not met, list number of days exceeded for each sample:			✓
11. Analysis Holding Time Met         If not met, list number of days exceeded for each sample:		✓	

# ADDITIONAL COMMENTS:

Recovery and RPD for some compounds are slightly biased high in PB167564BS/BSD but their hit is not present in any associated samples. The data will be used for hardcopies.



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A NO YES